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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
10/601,091	06/19/2003	Hong Xin	PP-01451.105	6161
27476	7590 12/15/2006		EXAMINER	
NOVARTIS VACCINES AND DIAGNOSTICS INC.			MARTINELL, JAMES	
CORPORATE INTELLECTUAL PROPERTY R338 P.O. BOX 8097			ART UNIT	PAPER NUMBER
Emeryville,	CA 94662-8097		1634	
		•	DATE MAILED: 12/15/2006	5

Please find below and/or attached an Office communication concerning this application or proceeding.

* ,	Application No.	Applicant(s)
	10/601,091	XIN ET AL.
Office Action Summary	Examiner	Art Unit
	James Martinell	1634
The MAILING DATE of this communication app Period for Reply	ears on the cover sheet with the c	orrespondence address
A SHORTENED STATUTORY PERIOD FOR REPLY WHICHEVER IS LONGER, FROM THE MAILING DA  - Extensions of time may be available under the provisions of 37 CFR 1.13 after SIX (6) MONTHS from the mailing date of this communication.  - If NO period for reply is specified above, the maximum statutory period w  - Failure to reply within the set or extended period for reply will, by statute, Any reply received by the Office later than three months after the mailing earned patent term adjustment. See 37 CFR 1.704(b).	ATE OF THIS COMMUNICATION 36(a). In no event, however, may a reply be tim vill apply and will expire SIX (6) MONTHS from cause the application to become ABANDONEI	l. ely filed the mailing date of this communication. D (35 U.S.C. § 133).
Status		
<ul> <li>1) Responsive to communication(s) filed on 05 Oc</li> <li>2a) This action is FINAL. 2b) This</li> <li>3) Since this application is in condition for allowar closed in accordance with the practice under E</li> </ul>	action is non-final. nce except for formal matters, pro	
Disposition of Claims		
4) ☑ Claim(s) 11-13,16,18-21,23,25,27,29,30,32,36, 4a) Of the above claim(s) 29,30,32,42,43,48 and 5) ☐ Claim(s) is/are allowed. 6) ☑ Claim(s) 11-13,16,18-21,23,25,27,36,39,40,44 7) ☐ Claim(s) is/are objected to. 8) ☐ Claim(s) are subject to restriction and/or	nd 49 is/are withdrawn from considered and 50-54 is/are rejected.	
Application Papers		
9)⊠ The specification is objected to by the Examine  10)⊠ The drawing(s) filed on 19 June 2003 is/are: a)  Applicant may not request that any objection to the Replacement drawing sheet(s) including the correct  11)□ The oath or declaration is objected to by the Ex	$\square$ accepted or b) $\square$ objected to drawing(s) be held in abeyance. See ion is required if the drawing(s) is obj	e 37 CFR 1.85(a). ected to. See 37 CFR 1.121(d).
Priority under 35 U.S.C. § 119		
a) All b) Some * c) None of:  1. Certified copies of the priority documents 2. Certified copies of the priority documents 3. Copies of the certified copies of the priority documents application from the International Bureau * See the attached detailed Office action for a list	s have been received. s have been received in Application ity documents have been received (PCT Rule 17.2(a)).	on No ed in this National Stage
Attachment(s)  1) Notice of References Cited (PTO-892)  2) Notice of Draftsperson's Patent Drawing Review (PTO-948)  3) Information Disclosure Statement(s) (PTO/SB/08) Paper No(s)/Mail Date 8/19/03 & 7/19/04	4) Interview Summary Paper No(s)/Mail Da 5) Notice of Informal P 6) Other:	ite

Applicant's election with traverse of the requirement for restriction in the reply filed on October 5, 2006 is acknowledged. The traversal is on the ground(s) that there is no undue burden of search to search Groups III and V and SEQ ID NOs: 1 and 11. This is not found persuasive because applicants have pointed to no structural relationship between SEQ ID NOs: 1 and 11. Searching is based on structural (*i.e.* sequence) relationships. Groups III and V are separate and distinct because Group V is concerned with measurement of gene expression products and not with the cDNA sequences *per se*.

The requirement is still deemed proper and is therefore made FINAL.

Claims 29, 30, 32, 42, 43, 48, and 49 are withdrawn from further consideration pursuant to 37 CFR 1.142(b), as being drawn to a nonelected invention, there being no allowable generic or linking claim. In addition, SEQ ID NO: 11 is withdrawn from consideration. Applicant timely traversed the restriction (election) requirement in the reply filed on October 5, 2006.

The drawings are objected to for the following reasons.

(a) The specification does not discuss Figures 3A, 3B, and 3C separately (*e.g.*, see page 6 of the application). This objection may be overcome by amendment to the description, drawings, or both.

Corrected drawing sheets in compliance with 37 CFR 1.121(d) are required in reply to the Office action to avoid abandonment of the application. Any amended replacement drawing sheet should include all of the figures appearing on the immediate prior version of the sheet, even if only one figure is being amended. The figure or figure number of an amended drawing should not be labeled as "amended." If a drawing figure is to be canceled, the appropriate figure must be removed from the replacement sheet, and where necessary, the remaining figures must be renumbered and appropriate changes made to the brief description of the several views of the drawings for consistency. Additional replacement sheets may be necessary to show the renumbering of the remaining figures. Each drawing sheet submitted after the filing date of an application must be labeled in the top margin as either "Replacement Sheet" or "New Sheet" pursuant to 37 CFR 1.121(d). If the changes are not accepted by the examiner, the applicant will

be notified and informed of any required corrective action in the next Office action. The objection to the drawings will not be held in abeyance.

The disclosure is objected to because of the following informalities.

- (a) A blank appears at page 1, line 3.
- (b) The instant application does not comply with the Sequence Rules (37 CFR §§ 1.821-1.825) because sequences appear without SEQ ID NO identifiers in at least the following locations, Figures 2A-1, 2A-2, 2B, 3A, 3B, and 3C. In order for any response to this Office action to be considered a complete response, the application must be amended to comply with the Sequence Rules. See MPEP 2422 and 37 CFR § 1.821(d).

Appropriate correction is required.

The following is a quotation of the first paragraph of 35 U.S.C. 112:

The specification shall contain a written description of the invention, and of the manner and process of making and using it, in such full, clear, concise, and exact terms as to enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make and use the same and shall set forth the best mode contemplated by the inventor of carrying out his invention.

The following is a quotation of the second paragraph of 35 U.S.C. 112:

The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.

Claim 25 is rejected under 35 U.S.C. 112, first paragraph, as failing to comply with the written description requirement. The claim(s) contains subject matter which was not described in the specification in such a way as to reasonably convey to one skilled in the relevant art that the inventor(s), at the time the application was filed, had possession of the claimed invention. The instant application does not adequately describe the genes metnioned in the claim. In *Vas-Cath v Mahurkar*, 19 USPQ2d 1111 (Fed. Cir. 1991) the court stated, "applicant must also convey with reasonable clarity to those skilled in the art that, as of the filing date sought, he or she was in possession *of the invention*. The invention is, for purposes of the 'written description' inquiry, *whatever is now claimed*' (emphasis in the

original) (*Vas-Cath* at page 1117). The instant application does not "clearly allow persons of ordinary skill in the art to recognize that [he or she] invented what is now claimed" (*Vas-Cath* at page 1116). In *Fiers v. Sugano*, 25 USPQ2d 1601 (Fed. Cir. 1993), the court also held that, "An adequate written description of a DNA requires more than a mere statement that it is part of the invention and reference a potential method for isolating it; what is required is a description of the DNA itself" (*Fiers v. Sugano*, page 1606). This view was reiterated in *Fiddes v. Baird*, USPQ2d 1481 (BPAI 1993) at page 1483, "If a conception of a DNA requires a specific definition, such as by structure, formula, chemical name, or physical properties, as we have held, then a description also requires that degree of specificity. . . . one cannot describe what one has not conceived." The court amplified this notion with respect to inventions claiming genetic material in *Regents of the University of California v. Eli Lilly*, 43 USPQ2d 1398 (Fed. Cir. 1997), stating at page 1406,

"In claims to genetic material, however, a generic statement such as 'vertebrate insulin cDNA' or mammalian insulin cDNA," without more, is not an adequate written description of the genus because it does not distinguish the claimed genus from others, except by function. It does not specifically define any of the genes that fall within its definition. It does not define any structural features commonly possessed by members of the genus that distinguish them from others. One skilled in the art therefore cannot, as one can do with a fully described genus, visualize or recognize the identity of the members of the genus. . . . Accordingly, naming a type of material generally known to exist, in the absence of knowledge as to what that material consists of, is not a description of that material."

Claims 11-13, 16, 18-21, 23, 25, 27, 36, 39, 40, 44-47, and 50-54 are rejected under 35 U.S.C. 112, second paragraph, as being indefinite for failing to particularly point out and distinctly claim the subject matter which applicant regards as the invention. The claims are vague, indefinite, and incomplete.

- (a) Claims 11-13, 16, 19, 21, 23, 25, 27, 29, 36, 39, 40, 44, and 46 claim more than was elected (*i.e.* SEQ ID NO: 11).
- (b) The recitation of "subgenomic polynucleotide" (claims 19 and 20) is vague and indefinite because the term is not defined and it has no clear art-recognized meaning.

(c) The recitation of "a gene" (claim 25) is vague and indefinite because the instant application does not clearly define the term and the term has no clear art-recognized single meaning.

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- (d) The recitation of "detectable label" (claim 27) is vague and indefinite because the instant application does not distinguish a detectable label from an undetectable label.
- (e) Claim 50 is incomplete because it depends from cancelled claim 1.

The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless -

- (b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on sale in this country, more than one year prior to the date of application for patent in the United States.
- (e) the invention was described in (1) an application for patent, published under section 122(b), by another filed in the United States before the invention by the applicant for patent or (2) a patent granted on an application for patent by another filed in the United States before the invention by the applicant for patent, except that an international application filed under the treaty defined in section 351(a) shall have the effects for purposes of this subsection of an application filed in the United States only if the international application designated the United States and was published under Article 21(2) of such treaty in the English language.

The following is a quotation of 35 U.S.C. 103(a) which forms the basis for all obviousness rejections set forth in this Office action:

(a) A patent may not be obtained though the invention is not identically disclosed or described as set forth in section 102 of this title, if the differences between the subject matter sought to be patented and the prior art are such that the subject matter as a whole would have been obvious at the time the invention was made to a person having ordinary skill in the art to which said subject matter pertains. Patentability shall not be negatived by the manner in which the invention was made.

This application currently names joint inventors. In considering patentability of the claims under 35 U.S.C. 103(a), the examiner presumes that the subject matter of the various claims was commonly owned at the time any inventions covered therein were made absent any evidence to the contrary.

Applicant is advised of the obligation under 37 CFR 1.56 to point out the inventor and invention dates of each claim that was not commonly owned at the time a later invention was made in order for the

RESULT 27 US-08-999-723-1

examiner to consider the applicability of 35 U.S.C. 103(c) and potential 35 U.S.C. 102(e), (f) or (g) prior art under 35 U.S.C. 103(a).

Claims 13, 16, 19, 20, 27, 36, and 44-46 are rejected under 35 U.S.C. 102(e) as being clearly anticipated by Powell et al (U.S. Patent No. 6,025,180). Powell et al discloses a nucleic acid that over its total length of 1862 nucleotides matches SEQ ID NO: 1 at 1400 positions (see the alignment below). Thus, the nucleic acid of Powell et al is embraced by the claims. Powell et al teaches PCR amplification (e.g., see column 10, line 52 through column 11, line 42 and column 16, lines 30-67), thus, any primers for the nucleic acid sequence of Powell et al are embraced by claim 36.

## ALIGNMENT OF SEQ ID NO: 1 WITH SEQ ID NO: 1 OF U.S. Patent No. 6,025,180

```
; Sequence 1, Application US/08999723A
; Patent No. 6025180
; GENERAL INFORMATION:
; APPLICANT: Powell, David J.
 APPLICANT: Southan, Christopher
 APPLICANT: Chapman, Conrad G.
  APPLICANT: Evans, Joanne R.
  TITLE OF INVENTION: ASP1
  FILE REFERENCE: GH70262
  CURRENT APPLICATION NUMBER: US/08/999,723A
  CURRENT FILING DATE: 1997-10-06
 NUMBER OF SEQ ID NOS: 2
  SOFTWARE: PatentIn Ver. 2.0
; SEO ID NO 1
   LENGTH: 1862
   TYPE: DNA
   ORGANISM: Homo sapiens
US-08-999-723-1
                       55.6%; Score 1350; DB 3; Length 1862;
 Query Match
 Best Local Similarity
                       99.9%; Pred. No. 0;
 Matches 1400; Conservative
                             0; Mismatches
                                             1:
                                               Indels
        268 AACCCCGCACTCCTACATAGACACGTACTTTGACACAGAGAGGTCTAGCACATACCGCTC 327
Qу
            450 AACCCCGCACTCCTACATAGACACGTACTTTGACACAGAGAGGTCTAGCACATACCGCTC 509
Db
        328 CAAGGGCTTTGACGTCACAGTGAAGTACACACAAGGAAGCTGGACGGCTTCGTTGGGGA 387
Qу
            510 CAAGGGCTTTGACGTCACAGTGAAGTACACACAAGGAAGCTGGACGGGCTTCGTTGGGGA 569
Db
        388 AGACCTCGTCACCATCCCCAAAGGCTTCAATACTTCTTTTCTTGTCAACATTGCCACTAT 447
Qу
```

Db	570	AGACCTCGTCACCATCCCCAAAGGCTTCAATACTTCTTTTCTTGTCAACATTGCCACTAT	629
Qу	448	TTTTGAATCAGAGAATTTCTTTTTGCCTGGGATTAAATGGAATGGAATACTTGGCCTAGC	507
Db	630	TTTTGAATCAGAGAATTTCTTTTTGCCTGGGATTAAATGGAATGGAATACTTGGCCTAGC	689
Qу	508	TTATGCCACACTTGCCAAGCCATCAAGTTCTCTGGAGACCTTCTTCGACTCCCTGGTGAC	567
Db	690	TTATGCCACACTTGCCAAGCCATCAAGTTCTCTGGAGACCTTCTTCGACTCCCTGGTGAC	749
Qу	568	ACAAGCAAACATCCCCAACGTTTTCTCCATGCAGATGTGTGGAGCCGGCTTGCCCGTTGC	627
Db	750 <sub>.</sub>	ACAAGCAAACATCCCCAACGTTTTCTCCATGCAGATGTGTGGAGCCGGCTTGCCCGTTGC	809
Qу	628	TGGATCTGGGACCAACGGAGGTAGTCTTGTCTTGGGTGGAATTGAACCAAGTTTGTATAA	687
Db	810	TGGATCTGGGACCAACGGAGGTAGTCTTGTCTTGGGTGGAATTGAACCAAGTTTGTATAA	869
QУ	688	AGGAGACATCTGGTATACCCCTATTAAGGAAGAGTGGTACTACCAGATAGAAATTCTGAA	747
Db	870	AGGAGACATCTGGTATACCCCTATTAAGGAAGAGTGGTACTACCAGATAGAAATTCTGAA	929
Qy	748	ATTGGAAATTGGAGGCCAAAGCCTTAATCTGGACTGCAGAGAGTATAACGCAGACAAGGC	807
Db	930	ATTGGAAATTGGAGGCCAAAGCCTTAATCTGGACTGCAGAGAGTATAACGCAGACAAGGC	989
Qу	808	CATCGTGGACAGTGGCACCACGCTGCTGCCCCAGAAGGTGTTTGATGCGGTGGT	867
Db	990	${\tt CATCGTGGACAGTGCCACGCTGCTGCCCCAGAAGGTGTTTGATGCGGTGGT}$	1049
Qу	868	GGAAGCTGTGGCCCGCGCATCTCTGATTCCAGAATTCTCTGATGGTTTCTGGACTGGGTC	927
Db	1050	GGAAGCTGTGGCCCGCGCATCTCTGATTCCAGAATTCTCTGATGGTTTCTGGACTGGGTC	1109
Qу	928	CCAGCTGGCGTGCTGGACGAATTCGGAAACACCTTGGTCTTACTTCCCTAAAATCTCCAT	987
Db	1110	$\tt CCAGCTGGCGTGCTGGACGAATTCGGAAACACCTTGGTCTTACTTCCCTAAAATCTCCAT$	1169
Qу	988	CTACCTGAGAGATGAGAACTCCAGCAGGTCATTCCGTATCACAATCCTGCCTCAGCTTTA	1047
Db	1170	CTACCTGAGAGACTCCAGCAGGTCATTCCGTATCACAATCCTGCCTCAGCTTTA	1229
Qy	1048	CATTCAGCCCATGATGGGGGCCGGCCTGAATTATGAATGTTACCGATTCGGCATTTCCCC	1107
Db	1230	CATTCAGCCCATGATGGGGGCCGGCCTGAATTATGAATGTTACCGATTCGGCATTTCCCC	1289
Qу	1108	ATCCACAAATGCGCTGGTGATCGGTGCCACGGTGATGGAGGGCTTCTACGTCATCTTCGA	1167
Db	1290	ATCCACAAATGCGCTGGTGATCGGTGCCACGGTGATGGAGGGCTTCTACGTCATCTTCGA	1349
Qy	1168	CAGAGCCCAGAAGAGGGTGGGCTTCGCAGCGAGCCCCTGTGCAGAAATTGCAGGTGCTGC	1227
Db	1350	CAGAGCCCAGAAGAGGGTGGGCTTCGCAGCGAGCCCCTGTGCAGAAATTGCAGGTGCTGC	1409
Qy	1228	AGTGTCTGAAATTTCCGGGCCTTTCTCAACAGAGGATGTAGCCAGCAACTGTGTCCCCGC	1287
Db	1410	AGTGTCTGAAATTTCCGGGCCTTTCTCAACAGAGGATGTAGCCAGCAACTGTGTCCCCGC	1469

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Qу
      1288 TCAGTCTTTGAGCGAGCCCATTTTGTGGATTGTGTCCTATGCGCTCATGAGCGTCTGTGG 1347
          Db
      1470 TCAGTCTTTGAGCGAGCCCATTTTGTGGATTGTGTCCTATGCGCTCATGAGCGTCTGTGG 1529
      1348 AGCCATCCTCCTTGTCTTAATCGTCCTGCTGCTGCCGTTCCGGTGTCAGCGTCGCCC 1407
Qу
          Db
      1530 AGCCATCCTCCTTGTCTTAATCGTCCTGCTGCTGCCGTTCCGGTGTCAGCGTCGCCC 1589
      1408 CCGTGACCCTGAGGTCGTCAATGATGAGTCCTCTCTGGTCAGACATCGCTGGAAATGAAT 1467
Qу
          Db
      1590 CCGTGACCCTGAGGTCGTCAATGATGAGTCCTCTCTGGTCAGACATCGCTGGAAATGAAT 1649
      1468 AGCCAGGCCTGACCTCAAGCAACCATGAACTCAGCTATTAAGAAAATCACATTTCCAGGG 1527
Qу
          Db
      1650 AGCCAGGCCTGACCTCAAGCAACCATGAACTCAGCTATTAAGAAAATCACATTTCCAGGG 1709
Qу
      1528 CAGCAGCCGGGATCGATGGTGGCGCTTTCTCCTGTGCCCACCCGTCTTCAATCTCTGTTC 1587
          Db
      1710 CAGCAGCCGGGATCGATGGTGGCGCTTTCTCCTGTGCCCACCCGTCTTCAATCTCTGTTC 1769
      1588 TGCTCCCAGATGCCTTCTAGATTCACTGTCTTTTGATTCTTGATTTTCAAGCTTTCAAAT 1647
Qу
          1770 TGCTCCCAGATGCCTTCTAGATTCACTGTCTTTTGATTCTTGATTTTCAAGCTTTCAAAT 1829
Db
      1648 CCTCCCTACTTCCAAGAAAA 1668
Qу
          Db
      1830 CCTCCCTACTTCCAAGAAAA 1850
```

Claims 13, 16, 19, 20, 27, and 44-46 are rejected under 35 U.S.C. 102(b) as being clearly anticipated by Hillier et al (GenBank® Accession No. AA056282 (November 29, 1996)). Hillier et al discloses a nucleic acid that over its total length of 618 nucleotides matches SEQ ID NO: 1 at 541 positions (see the alignment below). Thus, the nucleic acid of Hillier et al is embraced by the claims.

```
RESULT 54
AA056282
                                                                 EST 29-NOV-1996
            AA056282
                                      618 bp
                                                mRNA
                                                        linear
LOCUS
            ze24b09.rl Soares fetal heart NbHH19W Homo sapiens cDNA clone
DEFINITION
            IMAGE: 359897 5', mRNA sequence.
ACCESSION
            AA056282
VERSION
            AA056282.1 GI:1548686
KEYWORDS
            EST.
SOURCE
            Homo sapiens (human)
  ORGANISM
            Homo sapiens
            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
            Mammalia; Eutheria; Euarchontoglires; Primates; Catarrhini;
            Hominidae; Homo.
REFERENCE
            1 (bases 1 to 618)
  AUTHORS
            Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M.,
```

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Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M.,
           Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F.,
          Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P. and
          Wilson, R.
 TITLE
          The WashU-Merck EST Project
 JOURNAL
          Unpublished (1995)
COMMENT
          Contact: Wilson RK
          Washington University School of Medicine
           4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
          Tel: 314 286 1800
           Fax: 314 286 1810
          Email: est@watson.wustl.edu
          This clone is available royalty-free through LLNL; contact the
           IMAGE Consortium (info@image.llnl.gov) for further information.
           Putative full length read
          The vector to vector length is 619
           Insert Length: 404
                             Std Error: 0.00
          Seq primer: -28M13 rev2 from Amersham
          High quality sequence stop: 448.
FEATURES
                   Location/Qualifiers
    source
                   1. .618
                   /organism="Homo sapiens"
                   /mol type="mRNA"
                   /db xref="GDB:1276441"
                   /db xref="taxon:9606"
                   /clone="IMAGE:359897"
                   /sex="unknown"
                   /dev stage="19 weeks" ·
                   /lab host="DH10B (ampicillin resistant)"
                   /clone_lib="Soares_fetal_heart_NbHH19W"
                   /note="Organ: heart; Vector: pT7T3D (Pharmacia) with a
                   modified polylinker; Site 1: Not I; Site 2: Eco RI; 1st
                   strand cDNA was primed with a Not I - oligo(dT) primer [5'
                   double-stranded cDNA was size selected, ligated to Eco RI
                   adapters (Pharmacia), digested with Not I and cloned into
                   the Not I and Eco RI sites of a modified pT7T3 vector
                   (Pharmacia). Library went through one round of
                   normalization to a Cot = 5. Library constructed by
                   M. Fatima Bonaldo. This library was constructed from the
                   same fetus as the fetal lung library, Soares fetal lung
                   NbHL19W."
ORIGIN
                       22.3%; Score 541; DB 1; Length 618;
 Query Match
 Best Local Similarity
                       100.0%; Pred. No. 3.4e-267;
Matches 541; Conservative
                              0; Mismatches
                                              0; Indels
                                                           0; Gaps
                                                                      0;
        1798 CTTCTCTCTAATCTCTGGAAAAATAAGTACATATAGTTGATAACCCCTCTTAGCTTAC 1857
Qy
             1 CTTCTCTCTAATCTCTGGAAAAATAAGTACATATAGTTGATAACCCCTCTTAGCTTAC 60
Db
        1858 AGGAAGCTTTTTGTATTAATTGCCTTTGAGGTTATTTTCCGCCAGACCTCAACCTGGGTC 1917
Qу
            61 AGGAAGCTTTTTGTATTAATTGCCTTTGAGGTTATTTTCCGCCAGACCTCAACCTGGGTC 120
Db
        1918 AAAGTGGTACAGGAAGGCTTGCAGTATGATGGCAGGAGAATCAGCCTGGGGCCTGGGGAT 1977
Qу
            121 AAAGTGGTACAGGAAGGCTTGCAGTATGATGGCAGGAGAATCAGCCTGGGGCCTGGGGAT 180
Db
```

```
1978 GTAACCAAGCTGTAECCTTGAGACCTGGAACCAGAGCCACAGGCCCCTTTTGTGGGTTTC 2037
Qу
       Db
    181 GTAACCAAGCTGTACCCTTGAGACCTGGAACCAGAGCCACAGGCCCCTTTTGTGGGTTTC 240
    2038 TCTGTGCTCTGAATGGGAGCCAGAATTCACTAGGAGGTCATCAACCGATGGTCCTCACAA 2097
Qу
       Db
    241 TCTGTGCTCTGAATGGGAGCCAGAATTCACTAGGAGGTCATCAACCGATGGTCCTCACAA 300
Qу
    Db
    Qу
    2158 TTTTGTACCCAATACTTATGTTGTATTGTTGGTGCGAAAGTAAAAACACTACCTCTTTTG 2217
       Db
     361 TTTTGTACCCAATACTTATGTTGTATTGTTGGTGCGAAAGTAAAAACACTACCTCTTTTG 420
Qу
    Db
     2278 CCCCTCACCCAAAAGAATTATCATCCCAACAGCCAAGACCCAACAGGTGCTGAACTGTGC 2337
Qу
       481 CCCCTCACCCAAAAGAATTATCATCCCAACAGCCAAGACCCAACAGGTGCTGAACTGTGC 540
Db
    2338 A 2338
Qу
     541 A 541
Db
```

Claim 36 is rejected under 35 U.S.C. 103(a) as being unpatentable over Hillier et al (GenBank<sup>®</sup> Accession No. AA056282 (November 29, 1996) in view of applicants' admitted state of the prior art (*e.g.*, page 43, lines 20-22 of the application). Hillier et al discloses a nucleic acid that over its total length of 618 nucleotides matches SEQ ID NO: 1 at 541 positions (see the alignment above). Applicants acknowledge PCR amplification to be old (*e.g.*, instant application at page 43, lines 20-22). It would have been obvious for one of ordinary skill in the art at the time the invention was made to have constructed primers to hybridize to any portions of the nucleic acid disclosed by Hillier et al in order to make multiple copies of the DNA bounded by the primers for further analysis or production purposes.

Claims 39 and 40 are rejected under 35 U.S.C. 103(a) as being unpatentable over either one of Powell et al (U.S. Patent No. 6,025,180) or Hillier et al (GenBank<sup>®</sup> Accession No. AA056282 (November 29, 1996) in view of applicants' admitted state of the prior art (*e.g.*, page 41, lines 11-23 of the application). Powell et al discloses a nucleic acid that over its total length of 1862 nucleotides matches SEQ ID NO: 1 at 1400 positions (see the alignment above). Hillier et al discloses a nucleic acid that over its total length of 618 nucleotides matches SEQ ID NO: 1 at 541 positions (see the alignment above).

Applicants acknowledge use of arrays for nucleic acid detection to be old (*e.g.*, instant application at page 41, lines 11-23). It would have been obvious for one of ordinary skill in the art at the time the invention was made to have constructed arrays containing the nucleic acids or fragments of the nucleic acids disclosed in either one of Powell et al or Hillier et al in order to detect the nucleic acids of either one of Powell et al or Hillier et al in a given sample.

Any inquiry concerning this communication or earlier communications from the examiner should be directed to James Martinell whose telephone number is (571) 272-0719.

The examiner works a flexible schedule and can be reached by phone and voice mail.

Alternatively, a request for a return telephone call may be e-mailed to <a href="mailto:james.martinell@uspto.gov">james.martinell@uspto.gov</a>. Since e-mail communications may not be secure, it is suggested that information in such requests be limited to name, phone number, and the best time to return the call.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Ram Shukla, can be reached on (571) 272-0735.

## **OFFICIAL FAX NUMBER**

The fax phone number for the organization where this application or proceeding is assigned is (571) 273-8300.. Any Official Communication to the USPTO should be faxed to this number.

Any inquiry of a general nature or relating to the status of this application or proceeding should be directed to (571) 272-0547.

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